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1646

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/507,968B

Input Set : A:\09507968 SEQ LIST.txt

Output Set: N:\CRF3\04122001\I507968B.raw

ENTERED

3 <110> APPLICANT: Yu et al.
 5 <120> TITLE OF INVENTION: Neutrokin-alpha and Neutrokin-alpha Splice Variants
 7 <130> FILE REFERENCE: PF343P3
 9 <140> CURRENT APPLICATION NUMBER: 09/507,968B
 10 <141> CURRENT FILING DATE: 2000-02-22
 12 <150> PRIOR APPLICATION NUMBER: 60/122,388
 13 <151> PRIOR FILING DATE: 1999-03-01
 15 <150> PRIOR APPLICATION NUMBER: 60/124,097
 16 <151> PRIOR FILING DATE: 1999-03-12
 18 <150> PRIOR APPLICATION NUMBER: 60/126,599
 19 <151> PRIOR FILING DATE: 1999-03-26
 21 <150> PRIOR APPLICATION NUMBER: 60/127,598
 22 <151> PRIOR FILING DATE: 1999-04-01
 24 <150> PRIOR APPLICATION NUMBER: 60/130,412
 25 <151> PRIOR FILING DATE: 1999-04-16
 27 <150> PRIOR APPLICATION NUMBER: 60/130,696
 28 <151> PRIOR FILING DATE: 1999-04-23
 30 <150> PRIOR APPLICATION NUMBER: 60/131,278
 31 <151> PRIOR FILING DATE: 1999-04-27
 33 <150> PRIOR APPLICATION NUMBER: 03/255,794
 34 <151> PRIOR FILING DATE: 1999-02-23
 36 <150> PRIOR APPLICATION NUMBER: 60/131,673
 37 <151> PRIOR FILING DATE: 1999-04-29
 39 <150> PRIOR APPLICATION NUMBER: 60/136,784
 40 <151> PRIOR FILING DATE: 1999-05-28
 42 <150> PRIOR APPLICATION NUMBER: 60/142,659
 43 <151> PRIOR FILING DATE: 1999-07-06
 45 <150> PRIOR APPLICATION NUMBER: 60/145,824
 46 <151> PRIOR FILING DATE: 1999-07-27
 48 <150> PRIOR APPLICATION NUMBER: 60/167,239
 49 <151> PRIOR FILING DATE: 1999-11-24
 51 <150> PRIOR APPLICATION NUMBER: 60/168,624
 52 <151> PRIOR FILING DATE: 1999-12-03
 54 <150> PRIOR APPLICATION NUMBER: 60/171,108
 55 <151> PRIOR FILING DATE: 1999-12-16
 57 <150> PRIOR APPLICATION NUMBER: 60/171,626
 58 <151> PRIOR FILING DATE: 1999-12-23
 60 <150> PRIOR APPLICATION NUMBER: 60/176,015
 61 <151> PRIOR FILING DATE: 2000-01-14
 64 <160> NUMBER OF SEQ ID NOS: 38
 66 <170> SOFTWARE: PatentIn Ver. 2.1
 68 <210> SEQ ID NO: 1
 69 <211> LENGTH: 1100
 70 <212> TYPE: DNA
 71 <213> ORGANISM: Homo sapiens
 73 <220> FEATURE:
 74 <221> NAME/KEY: CDS

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75 <222> LOCATION: (147)..(1001)

77 <400> SEQUENCE: 1

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80 atcaacaaac acagataaca ggaaatgac cattcctgt ggtcattat tctaaaggcc 120
82 ccaacttcca aagttcaagt agtgat atg gat gac tcc aca gaa agg gag gag 173
83                                     Met Asp Asp Ser Thr Glu Arg Glu Gln
84                                     1           5
86 tca cgc ctt act tct tgc ctt aag aaa aga gaa gaa atg aaa ctg aag 221
87 Ser Arg Leu Thr Ser Cys Leu Lys Lys Arg Glu Glu Met Lys Leu Lys
88 10           15           20           25
90 gag tgt gtt tcc atc ctg cca cgg aag gaa agc ccc tct gtc cga tcc 269
91 Glu Cys Val Ser Ile Leu Pro Arg Lys Glu Ser Pro Ser Val Arg Ser
92           30           35           40
94 tcc aaa gac gga aag ctg ctg gct gca acc ttg ctg ctg gca ctg ctg 317
95 Ser Lys Asp Gly Lys Leu Leu Ala Ala Thr Leu Leu Leu Ala Leu Leu
96           45           50           55
98 tct tgc tgc ctg acg gtg gtg tct ttc tac cag gtg gcc gcc ctg caa 365
99 Ser Cys Cys Leu Thr Val Val Ser Phe Tyr Gln Val Ala Ala Leu Gln
100           60           65           70
102 ggg gac ctg gcc agc ctg cgg gca gag ctg cag gcc cac cag gag gag 413
103 Gly Asp Leu Ala Ser Leu Arg Ala Glu Leu Gln Gly His His Ala Glu
104           75           80           85
106 aag ctg cca gca gga gca gga gcc ccc aag gcc gcc ctg gag gaa gct 461
107 Lys Leu Pro Ala Gly Ala Gly Ala Pro Lys Ala Gly Leu Glu Glu Ala
108 90           95           100           105
110 cca gct gtc acc ggc gga ctg aaa atc ttt gaa cca cca gct cca gga 509
111 Pro Ala Val Thr Ala Gly Leu Lys Ile Phe Glu Pro Pro Ala Pro Gly
112           110           115           120
114 gaa ggc aac tcc agt cag aac agc aga aat aag cgt gcc gtt cag ggt 557
115 Glu Gly Asn Ser Ser Gln Asn Ser Arg Asn Lys Arg Ala Val Gln Gly
116           125           130           135
118 cca gaa gaa aca gtc act cca gac tgc ttg cca ctg att gca gac agt 605
119 Pro Glu Glu Thr Val Thr Gln Asp Cys Leu Gln Leu Ile Ala Asp Ser
120           140           145           150
122 gaa aca cca act ata cca aaa gga tct tac aca ttt gtt cca tgg ctt 653
123 Glu Thr Pro Thr Ile Gln Lys Gly Ser Tyr Thr Phe Val Pro Trp Leu
124           155           160           165
126 ctg agc ttt aaa agg gga agt gcc cta gaa gaa aaa gag aat aaa ata 701
127 Leu Ser Phe Lys Arg Gly Ser Ala Leu Glu Glu Lys Glu Asn Lys Ile
128 170           175           180           185
130 ttg gtc aaa gaa act ggt tac ttt ttt ata tat ggt cag gtt tta tat 749
131 Leu Val Lys Glu Thr Gly Tyr Phe Phe Ile Tyr Gly Gln Val Leu Tyr
132           190           195           200
134 act gat aag acc tac gcc atg gga cat cta att cag agg aag aag gtc 797
135 Thr Asp Lys Thr Tyr Ala Met Gly His Leu Ile Gln Arg Lys Lys Val
136           205           210           215
138 cat gtc ttt ggg gat gaa ttg agt ctg gtg act ttg ttt cga tgt att 845
139 His Val Phe Gly Asp Glu Leu Ser Leu Val Thr Leu Phe Arg Cys Ile
140           220           225           230

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142  aac aat atg cct gaa aca cta ccc aat aat tcc tgc tat tca gct ggc      893
143  gln asn met pro glu thr leu pro asn asn ser cys tyr ser ala gly
144      235                240                245
146  att aca aaa ctg gaa gaa gga gat gaa ctc caa ctt gca ata cca aga      941
147  ile ala lys leu glu glu gly asp glu leu gln leu ala ile pro arg
148  250                255                260                265
150  gaa aat gca caa ata tca ctg gat gga gat gtc aca ttt ttt ggt gca      989
151  glu asn ala gln ile ser leu asp gly asp val thr phe phe gly ala
152      270                275                280
154  tgg aaa ctg ctg tgacctactt acaccatgtg tgtagctatt ttctccctt      1041
155  leu lys leu leu
156      285
158  tctcgtgaac tataagaaga aagaatctaa ctgaaaatac caaaaaaaaa aaaaaaaaaa 1100
161  <110> SEQ ID NO: 2
162  <111> LENGTH: 285
163  <112> TYPE: PRT
164  <113> ORGANISM Homo sapiens
166  <400> SEQUENCE: 2
167  Met Asp Asp Ser Thr Glu Arg Glu Gln Ser Arg Leu Thr Ser Cys Leu
168      1          5          10          15
170  Lys Lys Arg Glu Glu Met Lys Leu Lys Glu Cys Val Ser Ile Leu Pro
171      20          25          30
173  Arg Lys Glu Ser Pro Ser Val Arg Ser Ser Lys Asp Gly Lys Leu Leu
174      35          40          45
176  Ala Ala Thr Leu Leu Leu Ala Leu Leu Ser Cys Cys Leu Thr Val Val
177      50          55          60
179  Ser Phe Tyr Gln Val Ala Ala Leu Gln Gly Asp Leu Ala Ser Leu Arg
180      65          70          75          80
182  Ala Glu Leu Gln Gly His His Ala Glu Lys Leu Pro Ala Gly Ala Gly
183      85          90          95
185  Ala Pro Lys Ala Gly Leu Glu Glu Ala Pro Ala Val Thr Ala Gly Leu
186      100         105         110
188  Lys Ile Phe Glu Pro Pro Ala Pro Gly Glu Gly Asn Ser Ser Gln Asn
189      115         120         125
191  Ser Arg Asn Lys Arg Ala Val Gln Gly Pro Glu Glu Thr Val Thr Gln
192      130         135         140
194  Asp Cys Leu Gln Leu Ile Ala Asp Ser Glu Thr Pro Thr Ile Gln Lys
195  145         150         155         160
197  Gly Ser Tyr Thr Phe Val Pro Trp Leu Leu Ser Phe Lys Arg Gly Ser
198      165         170         175
200  Ala Leu Glu Glu Lys Glu Asn Lys Ile Leu Val Lys Glu Thr Gly Tyr
201      180         185         190
203  Phe Phe Ile Tyr Gly Gln Val Leu Tyr Thr Asp Lys Thr Tyr Ala Met
204      195         200         205
206  Gly His Leu Ile Gln Arg Lys Lys Val His Val Phe Gly Asp Glu Leu
207      210         215         220
209  Ser Leu Val Thr Leu Phe Arg Cys Ile Gln Asn Met Pro Glu Thr Leu
210  225         230         235         240
212  Pro Asn Asn Ser Cys Tyr Ser Ala Gly Ile Ala Lys Leu Glu Glu Gly

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/507,968B

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Input Set : A:\09507968 SEQ LIST.txt

Output Set: N:\CRF3\04122001\I507968B.raw

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213          245          250          255
215 Asp Glu Leu Gln Leu Ala Ile Pro Arg Glu Asn Ala Gln Ile Ser Leu
216          260          265          270
218 Asp Gly Asp Val Thr Phe Phe Gly Ala Leu Lys Leu Leu
219          275          280          285
223 <210> SEQ ID NO: 3
224 <211> LENGTH: 233
225 <212> TYPE: PRT
226 <213> ORGANISM Homo sapiens
228 <400> SEQUENCE 3
229 Met Ser Thr Glu Ser Met Ile Arg Asp Val Glu Leu Ala Glu Glu Ala
230 1 5 10 15
232 Leu Pro Lys Lys Thr Gly Gly Pro Gln Gly Ser Arg Arg Cys Leu Phe
233 20 25 30
235 Leu Ser Leu Phe Ser Phe Leu Ile Val Ala Gly Ala Thr Thr Leu Phe
236 35 40 45
238 Cys Leu Leu His Phe Gly Val Ile Gly Pro Gln Arg Glu Glu Phe Pro
239 50 55 60
241 Arg Asp Leu Ser Leu Ile Ser Pro Leu Ala Gln Ala Val Arg Ser Ser
242 65 70 75 80
244 Ser Arg Thr Pro Ser Asp Lys Pro Val Ala His Val Val Ala Asn Pro
245 85 90 95
247 Gln Ala Glu Gly Gln Leu Gln Trp Leu Asn Arg Arg Ala Asn Ala Leu
248 100 105 110
250 Leu Ala Asn Gly Val Glu Leu Arg Asp Asn Gln Leu Val Val Pro Ser
251 115 120 125
253 Glu Gly Leu Tyr Leu Ile Tyr Ser Gln Val Leu Phe Lys Gly Gln Gly
254 130 135 140
256 Cys Pro Ser Thr His Val Leu Leu Thr His Thr Ile Ser Arg Ile Ala
257 145 150 155 160
259 Val Ser Tyr Gln Thr Lys Val Asn Leu Leu Ser Ala Ile Lys Ser Pro
260 165 170 175
262 Cys Gln Arg Glu Thr Pro Glu Gly Ala Glu Ala Lys Pro Trp Tyr Glu
263 180 185 190
265 Pro Ile Tyr Leu Gly Gly Val Phe Gln Leu Glu Lys Gly Asp Arg Leu
266 195 200 205
268 Ser Ala Glu Ile Asn Arg Pro Asp Tyr Leu Asp Phe Ala Glu Ser Gly
269 210 215 220
271 Gln Val Tyr Phe Gly Ile Ile Ala Leu
272 225 230
275 <210> SEQ ID NO: 4
276 <211> LENGTH: 205
277 <212> TYPE: PRT
278 <213> ORGANISM Homo sapiens
280 <400> SEQUENCE: 4
281 Met Thr Pro Pro Glu Arg Leu Phe Leu Pro Arg Val Arg Gly Thr Thr
282 1 5 10 15
284 Leu His Leu Leu Leu Leu Gly Leu Leu Leu Val Leu Leu Pro Gly Ala
285 20 25 30

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287 Gln Gly Leu Pro Gly Val Gly Leu Thr Pro Ser Ala Ala Gln Thr Ala
288      35      40      45
290 Arg Gln His Pro Lys Met His Leu Ala His Ser Thr Leu Lys Pro Ala
291      50      55      60
293 Ala His Leu Ile Gly Asp Pro Ser Lys Gln Asn Ser Leu Leu Trp Arg
294      65      70      75      80
296 Ala Asn Thr Asp Arg Ala Phe Leu Gln Asp Gly Phe Ser Leu Ser Asn
297      85      90      95
300 Asn Ser Leu Leu Val Pro Thr Ser Gly Ile Tyr Phe Val Tyr Ser Gln
301      100      105      110
302 Val Val Phe Ser Gly Lys Ala Tyr Ser Pro Lys Ala Thr Ser Ser Pro
303      115      120      125
305 Leu Tyr Leu Ala His Glu Val Gln Leu Phe Ser Ser Gln Tyr Pro Phe
306      130      135      140
308 His Val Pro Leu Leu Ser Ser Gln Lys Met Val Tyr Pro Gly Leu Gln
309      145      150      155      160
311 Glu Pro Trp Leu His Ser Met Tyr His Gly Ala Ala Phe Gln Leu Thr
312      165      170      175
314 Gln Gly Asp Gln Leu Ser Thr His Thr Asp Gly Ile Pro His Leu Val
315      180      185      190
317 Leu Ser Pro Ser Thr Val Phe Phe Gly Ala Phe Ala Leu
318      195      200      205
321 <210> SEQ ID NO: 5
322 <211> LENGTH: 244
323 <212> TYPE: PRT
324 <213> ORGANISM Homo sapiens
326 <400> SEQUENCE 5
327 Met Gly Ala Leu Gly Leu Glu Gly Arg Gly Gly Arg Leu Gln Gly Arg
328      1      5      10      15
330 Gly Ser Leu Leu Leu Ala Val Ala Gly Ala Thr Ser Leu Val Thr Leu
331      20      25      30
333 Leu Leu Ala Val Pro Ile Thr Val Leu Ala Val Leu Ala Leu Val Pro
334      35      40      45
336 Gln Asp Gln Gly Gly Leu Val Thr Glu Thr Ala Asp Pro Gly Ala Gln
337      50      55      60
339 Ala Gln Gln Gly Leu Gly Phe Gln Lys Leu Pro Glu Gln Glu Pro Glu
340      65      70      75      80
342 Thr Asp Leu Ser Pro Gly Leu Pro Ala Ala His Leu Ile Gly Ala Pro
343      85      90      95
345 Leu Lys Gly Gln Gly Leu Gly Trp Glu Thr Thr Lys Glu Gln Ala Phe
346      100      105      110
348 Leu Thr Ser Gly Thr Gln Phe Ser Asp Ala Glu Gly Leu Ala Leu Pro
349      115      120      125
351 Gln Asp Gly Leu Tyr Tyr Leu Tyr Cys Leu Val Gly Tyr Arg Gly Arg
352      130      135      140
354 Ala Pro Pro Gly Gly Gly Asp Pro Gln Gly Arg Ser Val Thr Leu Arg
355      145      150      155      160
357 Ser Ser Leu Tyr Arg Ala Gly Gly Ala Tyr Gly Pro Gly Thr Pro Glu
358      165      170      175

```

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/507,968B

DATE: 04/13/2001

TIME: 13:20:12

Input Set : A:\09507968 SEQ LIST.txt

Output Set: N:\CRF3\04122001\I507968B.raw

L:496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:497 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:499 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:500 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:501 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:684 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:687 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:690 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:691 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:692 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:783 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:784 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:785 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:786 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:787 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:788 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:789 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:1382 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:1406 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36